16239 `#12`

OIPE

RAW SEQUENCE LISTING   DATE: 07/05/2002
PATENT APPLICATION: US/09/890,811B   TIME: 15:37:23

Input Set : N:\Crf3\06122002\I890811A.raw
Output Set: N:\CRF3\07052002\I890811B.raw

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 1 <110> APPLICANT: Zhu, Qun
 2       Famodu, Omolayo O.
 3 <120> TITLE OF INVENTION: SPF1-Related Transcription Factors
 4 <130> FILE REFERENCE: BB1436 PCT
 5 <140> CURRENT APPLICATION NUMBER: US/09/890,811B
 6 <141> CURRENT FILING DATE: 2002-05-07
 7 <150> PRIOR APPLICATION NUMBER: 60/174325
 8 <151> PRIOR FILING DATE: 2000-01-04
 9 <160> NUMBER OF SEQ ID NOS: 14
10 <170> SOFTWARE: Microsoft Office 97
12 <210> SEQ ID NO: 1
13 <211> LENGTH: 512
14 <212> TYPE: DNA
15 <213> ORGANISM: Zea mays
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17 <221> NAME/KEY: unsure
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21 <221> NAME/KEY: unsure
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26 <222> LOCATION: (405)
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30 <222> LOCATION: (408)
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32 <220> FEATURE:
33 <221> NAME/KEY: unsure
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36 <220> FEATURE:
37 <221> NAME/KEY: unsure
38 <222> LOCATION: (447)
39 <223> OTHER INFORMATION: n=a,c,g or t
40 <220> FEATURE:
41 <221> NAME/KEY: unsure
42 <222> LOCATION: (453)
43 <223> OTHER INFORMATION: n=a,c,g or t
44 <220> FEATURE:
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C--> (line 5)

ENTERED

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    49 <221> NAME/KEY: unsure
    50 <222> LOCATION: (489)..(491)
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    52 <220> FEATURE:
    53 <221> NAME/KEY: unsure
    54 <222> LOCATION: (500)
    55 <223> OTHER INFORMATION: n=a,c,g or t
    56 <400> SEQUENCE: 1
    57     ctataattgg cgcaaatatg gacagaagca tgtcaaggga agtgaaaatc ctagaagtta   60
    58     ttacaagtgc actcatccta attgtgaagt taaaaagcta ttagagcgtt cgcttgatgg  120
    59     tcagattact gaagttgttt ataaagggca tcataatcat cccaagcccc aaccaaaatag 180
    60     aaggttagct gctggtgcag ttccttcaag ccaggctgaa gaaagatacg atggtgtggc  240
    61     acctattgaa gacaaacctt caaatattta ttccaacctc tgtaaccaag cacattcagc  300
    62     tggcatggtt gataatgttc cgggtccagc aagtgatgat gatgtttgatg ctggaggtgg  360
W--> 63     aagacccnac cctgggggga tgactcnaat gatgatgatg acttnggnct caaaaacgca  420
W--> 64     aggaaaatgg aatctgccgg gtatncnatg ccnggcttg antggggtaa accaaatccc  480
W--> 65     ggagccccnn nttcctttn aaaaactgtt tt                                  512
    67 <210> SEQ ID NO: 2
    68 <211> LENGTH: 170
    69 <212> TYPE: PRT
    70 <213> ORGANISM: Zea mays
    71 <220> FEATURE:
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    73 <222> LOCATION: (123)
    74 <223> OTHER INFORMATION: Xaa = ANY AMINO ACID
    75 <220> FEATURE:
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    77 <222> LOCATION: (129)
    78 <223> OTHER INFORMATION: Xaa = ANY AMINO ACID
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    81 <222> LOCATION: (135)..(136)
    82 <223> OTHER INFORMATION: Xaa = ANY AMINO ACID
    83 <220> FEATURE:
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    85 <222> LOCATION: (148)..(149)
    86 <223> OTHER INFORMATION: Xaa = ANY AMINO ACID
    87 <220> FEATURE:
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    94 <223> OTHER INFORMATION: Xaa = ANY AMINO ACID
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 96 <221> NAME/KEY: unsure
 97 <222> LOCATION: (163)..(164)
 98 <223> OTHER INFORMATION: Xaa = ANY AMINO ACID
 99 <220> FEATURE:
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101 <222> LOCATION: (167)
102 <223> OTHER INFORMATION: Xaa = ANY AMINO ACID
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104      Tyr Asn Trp Arg Lys Tyr Gly Gln Lys His Val Lys Gly Ser Glu Asn
105       1               5                  10                  15
106      Pro Arg Ser Tyr Tyr Lys Cys Thr His Pro Asn Cys Glu Val Lys Lys
107                      20                  25                  30
108      Leu Leu Glu Arg Ser Leu Asp Gly Gln Ile Thr Glu Val Val Tyr Lys
109                  35                  40                  45
110      Gly His His Asn His Pro Lys Pro Gln Pro Asn Arg Arg Leu Ala Ala
111              50                  55                  60
112      Gly Ala Val Pro Ser Ser Gln Ala Glu Glu Arg Tyr Asp Gly Val Ala
113      65                  70                  75                  80
114      Pro Ile Glu Asp Lys Pro Ser Asn Ile Tyr Ser Asn Leu Cys Asn Gln
115                      85                  90                  95
116      Ala His Ser Ala Gly Met Val Asp Asn Val Pro Gly Pro Ala Ser Asp
117                 100                 105                 110
118      Asp Asp Val Asp Ala Gly Gly Gly Arg Pro Xaa Pro Gly Gly Met Thr
119                 115                 120                 125
120      Xaa Met Met Met Met Thr Xaa Xaa Ser Lys Thr Gln Gly Lys Trp Asn
121                 130                 135                 140
122      Leu Pro Gly Xaa Xaa Cys Xaa Ala Leu Xaa Gly Val Asn Gln Ile Pro
123      145                 150                 155                 160
124      Glu Pro Xaa Xaa Pro Phe Xaa Lys Leu Phe
125                     165                 170
126 <210> SEQ ID NO: 3
127 <211> LENGTH: 717
128 <212> TYPE: DNA
129 <213> ORGANISM: Zea mays
130 <220> FEATURE:
131 <221> NAME/KEY: unsure
132 <222> LOCATION: (11)
133 <223> OTHER INFORMATION: n=a,c,g or t
134 <220> FEATURE:
135 <221> NAME/KEY: unsure
136 <222> LOCATION: (15)
137 <223> OTHER INFORMATION: n=a,c,g or t
138 <220> FEATURE:
139 <221> NAME/KEY: unsure
140 <222> LOCATION: (610)
141 <223> OTHER INFORMATION: n=a,c,g or t
142 <220> FEATURE:
143 <221> NAME/KEY: unsure
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W--> 118
W--> 120
W--> 122
W--> 124

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145 <222> LOCATION: (640)
146 <223> OTHER INFORMATION: n=a,c,g or t
147 <220> FEATURE:
148 <221> NAME/KEY: unsure
149 <222> LOCATION: (653)
150 <223> OTHER INFORMATION: n=a,c,g or t
151 <220> FEATURE:
152 <221> NAME/KEY: unsure
153 <222> LOCATION: (680)
154 <223> OTHER INFORMATION: n=a,c,g or t
155 <220> FEATURE:
156 <221> NAME/KEY: unsure
157 <222> LOCATION: (692)
158 <223> OTHER INFORMATION: n=a,c,g or t
159 <220> FEATURE:
160 <221> NAME/KEY: unsure
161 <222> LOCATION: (703)
162 <223> OTHER INFORMATION: n=a,c,g or t
163 <400> SEQUENCE: 3
164     gacgcacccc ntctnttctc tccccctctc gtctcgtcca gtcgtccccc tctcccccac   60
165     tctacccagc tgccctgctc tgcgtcgctc gccatggcgt cctccacggg gagcttggag  120
166     cacgggggt tcacgttcac gccgccgccc ttcatcacct ccttcaccga gctgctctcc  180
167     ggcgcagggg acatgctagg agccggcgcc gatcaggagc ggtcgtcgcc gaggggggctg  240
168     ttccaccgcg gcgccagggg cgtgcccaag ttcaagtcgg cgcagcctcc cagcctgccc  300
169     atctcgccgc cgcccatgtc gccttcctcc tacttcgcca tcccgcccgg gtcagcccc  360
170     gccgagctgc tcgactcgcc cgtcctgctc cactcgtcct ccaacatcct ggcgtctccc  420
171     accactggcg ccatcccggc gcagaggttc gactggaaga aggccgccga cctgatcgcg  480
172     tctcagtctc agcaagacgg cgacagccgg gctgccgccg ccggcttcga cgacttctcc  540
173     ttcacacggg caccttcaac gccgtgcgcg cgcacacgac gacgacgtcc ttaccttcat  600
174     gaagaagaan gtggagaggg ccctggccga cgggcgcatn aacgcaaaat cgngtacaaa  660
175     ggcgcggcac aaacaacccn aagccggttg tncacgcgcc ggnaacttaa ttccgga     717
177 <210> SEQ ID NO: 4
178 <211> LENGTH: 150
179 <212> TYPE: PRT
180 <213> ORGANISM: Zea mays
181 <400> SEQUENCE: 4
182     Met Ala Ser Ser Thr Gly Ser Leu Glu His Gly Gly Phe Thr Phe Thr
183      1               5                  10                  15
184     Pro Pro Pro Phe Ile Thr Ser Phe Thr Glu Leu Leu Ser Gly Ala Gly
185                  20                  25                  30
186     Asp Met Leu Gly Ala Gly Ala Asp Gln Glu Arg Ser Ser Pro Arg Gly
187                  35                  40                  45
188     Leu Phe His Arg Gly Ala Arg Gly Val Pro Lys Phe Lys Ser Ala Gln
189              50                  55                  60
190     Pro Pro Ser Leu Pro Ile Ser Pro Pro Pro Met Ser Pro Ser Ser Tyr
191      65                  70                  75                  80
192     Phe Ala Ile Pro Pro Gly Leu Ser Pro Ala Glu Leu Leu Asp Ser Pro
193                  85                  90                  95
194     Val Leu Leu His Ser Ser Ser Asn Ile Leu Ala Ser Pro Thr Thr Gly
```

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                    RAW SEQUENCE LISTING            DATE: 07/05/2002
                    PATENT APPLICATION:  US/09/890,811B    TIME: 15:37:23

                    Input Set : N:\Crf3\06122002\I890811A.raw
                    Output Set: N:\CRF3\07052002\I890811B.raw

195                      100                  105                  110
196        Ala Ile Pro Ala Gln Arg Phe Asp Trp Lys Lys Ala Ala Asp Leu Ile
197             115                  120                  125
198        Ala Ser Gln Ser Gln Gln Asp Gly Asp Ser Arg Ala Ala Ala Ala Gly
199             130                  135                  140
200        Phe Asp Asp Phe Ser Phe
201             145                  150
203 <210> SEQ ID NO: 5
204 <211> LENGTH: 1961
205 <212> TYPE: DNA
206 <213> ORGANISM: Oryza sativa
207 <400> SEQUENCE: 5
208        agtcgtctcg ttctcgtctc cgatcactct cctcctcatc ttcgtcacgg tctcctcgct    60
209        tcgctagctc gcttgcttgc tggctgagct gtggtacgct cgccatggcg tcctcgacgg   120
210        gggggttgga ccacgggttc acgttccgc cgccgccgtt catcacgtcg ttcaccgagc   180
211        tgctgtcggg gggcggtggg gacctgctcg gcgccggcgg tgaggagcgc tcgccgaggg   240
212        ggttctccag aggcggagcg agggtgggcg gcggggtgcc caagttcaag tccgcgcagc   300
213        cgccgagcct gccgctctcg ccgccgcccg tgtcgccgtc gtcctacttc gccatcccgc   360
214        cggggctcag cccaccgag ctgctcgact ccccgtcct cctcagctcc tcccatatct   420
215        tggcgttccc gaccaccggt gcaatcccgg ctcagaggta cgactggaag gccagcgccg   480
216        atctcatcgc ttctcagcaa gatgacagcc gcggcgactt ctccttccac accaactccg   540
217        acgccatggc cgcgcaaccg gcctcttcc cttccttcaa ggagcaagag cagcaagtgg   600
218        tcgagtcgag caagaacggc gccgccgccg cgtcgagcaa caagagcggc ggcggcggga   660
219        acaacaagct ggaggacggg tacaactgga ggaagtacgg gcagaagcag gtgaagggga   720
220        gcgagaaccc gaggagctac tacaagtgca cctacaacgg ctgctccatg aagaagaagg   780
221        tggagcgctc gctcgccgac ggccgcatca cccagatcgt ctacaagggc gcacacaacc   840
222        acccaagccc gctctccacc gccgcaacgc ctcttccggc tccaccgccg ccgcctgccg   900
223        cgacgacctc gcggcgcccg gcgcgggcgc ggaccagtac tccgccgcga cgcccgagaa   960
224        ctcctccgtc acgttcggcg acgacgaggc cgacaacgca tcgcaccgca gcgagggcga  1020
225        cgagcccgaa gccaagcgct ggaaaggagg atgctgacaa cgagggcagc tccggcggca  1080
226        tgggcggcgg cgccggcggc aacccggtgc gcgagccgag gcttgtggtg cagacgctga  1140
227        gcgacatcga catcctcgac aacggcttcc ggtggaggaa gtacggccag aagtcgtca  1200
228        agggcaaccc caacccaagg agctactaca agtcacgac ggtgggctgc ccggtgcgga  1260
229        agcacgtgga gcgggcgtcg cacgacacgc cacgacctgat caccacctac gagggcaagc  1320
230        acaaccacga cgtcccggta cggccgcggc ggcggcggcg gacgcgcccc ggcgccggcg  1380
231        tcgcctacgg ctggggccga tccgggccga cggacgtcgc cgccgcccag caggggccct  1440
232        acaccctcga gatgctcccc aaccccgccg gcctctacgg cggctacggc gccggcgccg  1500
233        gcggcgccgc gttcccgcgc accaaggacg agcggcggga cgacctgttc gtcgagtcgc  1560
234        tcctctgcta gtcgagccga gccgagccga gctgagctgg gccccacatc cccctgctcg  1620
235        ccacgtggcg tattttcgcc tcgccgtata cgtacggccg tatagcgtac gtatacacgc  1680
236        tcgcacgccc tgcccaacac ggcaatacac acatacatac tctcgtacac acgtagtagc  1740
237        atacatatac agtatagtag gtgtgtag tgg tagctagcta gggagtgaga tccaatttgt  1800
238        tgattcgttg caggccactg ccacgtgggc cacaccggaa acagtacacg cgtatacacc  1860
239        acacttggga tacgcgtacg tacgcacatg tacacgtagt tttgtgcctt tgtaactgct  1920
240        gagagacagg tcaaataaga ctgatgaatt tttcatttct t                      1961
242 <210> SEQ ID NO: 6
243 <211> LENGTH: 488
244 <212> TYPE: PRT
245 <213> ORGANISM: Oryza sativa
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RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 07/05/2002
PATENT APPLICATION: US/09/890,811B   TIME: 15:37:24

Input Set : N:\Crf3\06122002\I890811A.raw
Output Set: N:\CRF3\07052002\I890811B.raw

Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
Sequence Listing to ensure that a corresponding explanation is presented in the <220>
to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; N Pos. 368,387,405,408,445,447,453,462,489,490,491,500
Seq#:2; Xaa Pos. 123,129,135,136,148,149,151,154,163,164,167
Seq#:3; N Pos. 11,15,610,640,653,680,692,703

Invalid Line Length:
The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:14; Line(s) 743

Input Set : **N:\Crf3\06122002\I890811A.raw**
Output Set: **N:\CRF3\07052002\I890811B.raw**

L:5 M:270 C: Current Application Number differs, Wrong Format
L:63 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:360
L:64 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:420
L:65 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:480
L:118 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:112
L:120 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:128
L:122 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:144
L:124 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:160
L:164 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0
L:174 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:600
L:175 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:660